SCORE Search Results Details for Application 10801292 and Search Result us-10-801-292-1.rnpbm.

This page gives you Search Results detail for the Application 10801292 and Search Result us-10-801-292-1.rnpbm.

start

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                      GenCore version 5.1.9
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OM nucleic - nucleic search, using sw model

Run on:         October 15, 2006, 07:51:11 ; Search time 2626 Seconds
                                           (without alignments)
                                           9807.648 Million cell updates/sec

Title:          US-10-801-292-1
Perfect score:  2096
Sequence:       1 ctcctctgctcctcgaagaa..........aaaaaaaaaaaaaaaaaaaa 2096

Scoring table:  IDENTITY_NUC
                Gapop 10.0 , Gapext 1.0

Searched:       18892170 seqs, 6143817638 residues

Total number of hits satisfying chosen parameters:      37784340

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

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        Pred. No. is the number of results predicted by chance to have a
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                              SUMMARIES
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Result          Query
  No.   Score   Match Length DB  ID                 Description
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	9	594	28.3	625	6	US-10-066-543-481	Sequence 481, App
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	45	157	7.5	996	3	US-09-876-143-1286	Sequence 1286, Ap

ALIGNMENTS

RESULT 1
US-10-801-292-1
; Sequence 1, Application US/10801292
; Publication No. US20050202448A1
; GENERAL INFORMATION:
; APPLICANT: LEE, YI-CHAO
; APPLICANT: YUEN, PUI-YEE
; APPLICANT: HUANG, YI-HUEI
; APPLICANT: WU, HUI-CHUAN
; TITLE OF INVENTION: METHODS FOR IDENTIFICATION, ASSESSMENT, PREVENTION, AND
; TITLE OF INVENTION: THERAPY OF CANCER
; FILE REFERENCE: 5422-2
; CURRENT APPLICATION NUMBER: US/10/801,292
; CURRENT FILING DATE: 2004-03-15
; NUMBER OF SEQ ID NOS: 21
; SOFTWARE: PatentIn Ver. 3.2
; SEQ ID NO 1
;   LENGTH: 2096
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-10-801-292-1

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  Matches 2096; Conservative   0; Mismatches   0; Indels   0; Gaps   0;

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US-10-756-149-4599
; Sequence 4599, Application US/10756149
; Publication No. US20050181375A1
; GENERAL INFORMATION:
;   APPLICANT: Aziz, Natasha
;   APPLICANT:  Zlotnik, Albert
;   TITLE OF INVENTION: NOVEL METHODS OF DIAGNOSIS OF METASTATIC CANCER, COMPOSITIONS AND
;   TITLE OF INVENTION:  METHODS OF SCREENING FOR MODULATORS OF METASTATIC CANCER
;   FILE REFERENCE: file
;   CURRENT APPLICATION NUMBER: US/10/756,149
;   CURRENT FILING DATE:  2004-01-12
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;  SOFTWARE: PatentIn version 3.2
; SEQ ID NO 4599
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;   TYPE: DNA
;   ORGANISM: Homo Sapiens
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (2508)..(2508)
;   OTHER INFORMATION: n is a, c, g, or t
US-10-756-149-4599

  Query Match          96.6%;  Score 2025.2;  DB 10;  Length 2527;
  Best Local Similarity  99.9%;  Pred. No. 0;
  Matches 2027;  Conservative   0;  Mismatches   3;  Indels   0;  Gaps   0;

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SCORE Search Results Details for Application 10801292 and Search Result us-10-801-292-1.rni.

This page gives you Search Results detail for the Application 10801292 and Search Result us-10-801-292-1.rni.

start

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                        GenCore version 5.1.9
              Copyright (c) 1993 - 2006  Biocceleration Ltd.


OM nucleic - nucleic search, using sw model

Run on:          October 15, 2006, 07:34:53 ; Search time 377 Seconds
                                     (without alignments)
                                     10402.769 Million cell updates/sec

Title:           US-10-801-292-1
Perfect score:   2096
Sequence:        1 ctcctctgctcctcgaagaa..........aaaaaaaaaaaaaaaaaaaa 2096

Scoring table:   IDENTITY_NUC
                 Gapop 10.0 , Gapext 1.0

Searched:        1403666 seqs, 935554401 residues

Total number of hits satisfying chosen parameters:      2807332

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
                 Maximum Match 100%
                 Listing first 45 summaries

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       Pred. No. is the number of results predicted by chance to have a
       score greater than or equal to the score of the result being printed,
       and is derived by analysis of the total score distribution.
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       2   259.6   12.4   2080  3   US-09-270-117-2        Sequence 2, Appli
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; Sequence 2, Application US/08878563A
; Patent No. 5891674
;  GENERAL INFORMATION:
;     APPLICANT:  Hillman, Jennifer L.
;     APPLICANT:  Lal, Preeti
;     APPLICANT:  Shah, Purvi
;     TITLE OF INVENTION:  INSULIN RECEPTOR TYROSINE KINASE SUBSTRATE
;     NUMBER OF SEQUENCES:  3
;     CORRESPONDENCE ADDRESS:
;       ADDRESSEE:  Incyte Pharmaceuticals, Inc.
;       STREET:  3174 Porter Drive
;       CITY:  Palo Alto
;       STATE:  CA
;       COUNTRY:  USA
;       ZIP:  94304
;     COMPUTER READABLE FORM:
;       MEDIUM TYPE:  Diskette
;       COMPUTER:  IBM Compatible
;       OPERATING SYSTEM:  DOS
;       SOFTWARE:  FastSEQ for Windows Version 2.0
;     CURRENT APPLICATION DATA:
;       APPLICATION NUMBER:  US/08/878,563A
;       FILING DATE:  Filed Herewith
;     PRIOR APPLICATION DATA:
;       APPLICATION NUMBER:
;       FILING DATE:
;     ATTORNEY/AGENT INFORMATION:
;       NAME:  Billings, Lucy J.
;       REGISTRATION NUMBER:  36,749
;       REFERENCE/DOCKET NUMBER:  PF-0323 US
;     TELECOMMUNICATION INFORMATION:
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;       TELEPHONE:  415-855-0555
;       TELEFAX:  415-845-4166
;  INFORMATION FOR SEQ ID NO:  2:
;     SEQUENCE CHARACTERISTICS:
;       LENGTH:  2080 base pairs
;       TYPE:  nucleic acid
;       STRANDEDNESS:  single
;       TOPOLOGY:  linear
;     IMMEDIATE SOURCE:
;       LIBRARY:  BRSTNOT04
;       CLONE:  918158
US-08-878-563A-2
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  Best Local Similarity  59.2%;  Pred. No. 1.6e-48;
  Matches  442;  Conservative    0;  Mismatches  304;  Indels    0;  Gaps    0;
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          I  II      II IIII    II      II III    IIIIIIIII I II
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; Patent No. 6265550
;   GENERAL INFORMATION:
;     APPLICANT:  Hillman, Jennifer L.
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;   APPLICANT: Lal, Preeti
;   APPLICANT: Shah, Purvi
;   TITLE OF INVENTION:  INSULIN RECEPTOR TYROSINE KINASE SUBSTRATE
;   NUMBER OF SEQUENCES:  3
;   CORRESPONDENCE ADDRESS:
;     ADDRESSEE:  Incyte Pharmaceuticals, Inc.
;     STREET:  3174 Porter Drive
;     CITY:  Palo Alto
;     STATE:  CA
;     COUNTRY:  USA
;     ZIP:  94304
;   COMPUTER READABLE FORM:
;     MEDIUM TYPE:  Diskette
;     COMPUTER:  IBM Compatible
;     OPERATING SYSTEM:  DOS
;     SOFTWARE:  FastSEQ for Windows Version 2.0
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;     APPLICATION NUMBER:  US/09/270,117
;     FILING DATE:
;   PRIOR APPLICATION DATA:
;     APPLICATION NUMBER:  08/878,563
;     FILING DATE:
;   ATTORNEY/AGENT INFORMATION:
;     NAME:  Billings, Lucy J.
;     REGISTRATION NUMBER:  36,749
;     REFERENCE/DOCKET NUMBER:  PF-0323 US
;   TELECOMMUNICATION INFORMATION:
;     TELEPHONE:  415-855-0555
;     TELEFAX:  415-845-4166
;  INFORMATION FOR SEQ ID NO:  2:
;   SEQUENCE CHARACTERISTICS:
;     LENGTH:  2080 base pairs
;     TYPE:  nucleic acid
;     STRANDEDNESS:  single
;     TOPOLOGY:  linear
;   IMMEDIATE SOURCE:
;     LIBRARY:  BRSTNOT04
;     CLONE:  918158
US-09-270-117-2
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Qy      590 AACACAAGAATAAATTAGAGTCTTTGGAGAAATCCCAAGCTGAGTTGAAGAAGATCAGAA 649
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;  GENERAL INFORMATION:
;    APPLICANT:  Hillman, Jennifer L.
;    APPLICANT:  Corley, Neil C.
;    APPLICANT:  Guegler, Karl J.
;    APPLICANT:  Baughn, Mariah
;    TITLE OF INVENTION:  INSULIN RECEPTOR TYROSINE KINASE SUBSTRATE
;    NUMBER OF SEQUENCES:  5
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Incyte Pharmaceuticals, Inc.
;      STREET:  3174 Porter Dr.
;      CITY:  Palo Alto
;      STATE:  CA
;      COUNTRY:  USA
;      ZIP:  94304
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Diskette
;      COMPUTER:  IBM Compatible
;      OPERATING SYSTEM:  DOS
;      SOFTWARE:  FastSEQ for Windows Version 2.0
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;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:
;      FILING DATE:
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Billings, Lucy J.
;      REGISTRATION NUMBER:  36,749
;      REFERENCE/DOCKET NUMBER:  PF-0323-1 CIP
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  650-855-0555
;      TELEFAX:  650-845-4166
; ·INFORMATION FOR SEQ ID NO:  2:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  2080 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    IMMEDIATE SOURCE:
;      LIBRARY:  BRSTNOT06
;      CLONE:  918158
US-09-046-572-2

   Query Match         12.4%;  Score 259.6;  DB 3;  Length 2080;
   Best Local Similarity  59.2%;  Pred. No. 1.6e-48;
   Matches  442;  Conservative    0;  Mismatches  304;  Indels    0;  Gaps    0;

Qy    230 CCGAGGAGGTGAACCGGCTCACGGAGAGCACCTACCGGAATGTTATGGAACAGTTCAATC 289
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                        GenCore version 5.1.9
              Copyright (c) 1993 - 2006  Biocceleration Ltd.


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      score greater than or equal to the score of the result being printed,
      and is derived by analysis of the total score distribution.

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	12	644.2	30.7	650	8	US-11-266-748A-15168	Sequence 15168, A
	13	630.8	30.1	634	8	US-11-266-748A-13471	Sequence 13471, A
	14	568.4	27.1	699	8	US-11-266-748A-367035	Sequence 367035,
c	15	568.4	27.1	699	8	US-11-266-748A-450414	Sequence 450414,
	16	505	24.1	506	8	US-11-266-748A-44783	Sequence 44783, A
	17	498.2	23.8	1000	8	US-11-266-748A-282338	Sequence 282338,
c	18	498.2	23.8	1000	8	US-11-266-748A-308978	Sequence 308978,
c	19	398.6	19.0	654	8	US-11-266-748A-212284	Sequence 212284,
	20	398.6	19.0	654	8	US-11-266-748A-235743	Sequence 235743,
c	21	389	18.6	1214	8	US-11-266-748A-254365	Sequence 254365,
	22	389	18.6	1214	8	US-11-266-748A-314882	Sequence 314882,
	23	319.4	15.2	869	8	US-11-266-748A-44784	Sequence 44784, A
c	24	308.4	14.7	447	8	US-11-266-748A-163280	Sequence 163280,
	25	308.4	14.7	447	8	US-11-266-748A-243773	Sequence 243773,
	26	258	12.3	1684	8	US-11-266-748A-261669	Sequence 261669,
	27	258	12.3	1684	8	US-11-266-748A-279563	Sequence 279563,
c	28	258	12.3	1684	8	US-11-266-748A-322186	Sequence 322186,
	29	258	12.3	2129	8	US-11-266-748A-22362	Sequence 22362, A
	30	258	12.3	2129	8	US-11-266-748A-29713	Sequence 29713, A
	31	258	12.3	2519	8	US-11-266-748A-26814	Sequence 26814, A
	32	256.4	12.2	1009	8	US-11-266-748A-93448	Sequence 93448, A
	33	256.4	12.2	1009	8	US-11-266-748A-113087	Sequence 113087,
c	34	256.4	12.2	1009	8	US-11-266-748A-146259	Sequence 146259,
c	35	253.2	12.1	609	8	US-11-266-748A-172311	Sequence 172311,
	36	253.2	12.1	609	8	US-11-266-748A-245546	Sequence 245546,
	37	241.2	11.5	924	8	US-11-266-748A-171204	Sequence 171204,
	38	99.2	4.7	826	8	US-11-266-748A-454	Sequence 454, App
	39	96.8	4.6	1000	8	US-11-266-748A-397933	Sequence 397933,
c	40	96.8	4.6	1000	8	US-11-266-748A-468979	Sequence 468979,
c	41	83.4	4.0	500	7	US-11-292-078-15999	Sequence 15999, A
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	43	82.8	4.0	997	6	US-10-196-749-23	Sequence 23, Appl
c	44	82.4	3.9	574	7	US-11-292-078-11147	Sequence 11147, A
	45	81.8	3.9	1000	8	US-11-266-748A-398495	Sequence 398495,

ALIGNMENTS

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RESULT 1
US-11-266-748A-227410
; Sequence 227410, Application US/11266748A
; Publication No. US20060134663A1
; GENERAL INFORMATION:
;  APPLICANT: Harkin, Paul
;  APPLICANT:  Johnston, Patrick
;  APPLICANT:  Mulligan, Karl
;  TITLE OF INVENTION: Transcriptome Microarray Technology and
;  TITLE OF INVENTION:  Methods of Using the Same
;  FILE REFERENCE: 55815-0102 (319189)
;  CURRENT APPLICATION NUMBER: US/11/266,748A
;  CURRENT FILING DATE:  2005-11-03
;  PRIOR APPLICATION NUMBER: EP 04105479.2
;  PRIOR FILING DATE: 2004-11-03
;  PRIOR APPLICATION NUMBER: EP 04105482.6
;  PRIOR FILING DATE: 2004-11-03
;  PRIOR APPLICATION NUMBER: EP 04105483.4
;  PRIOR FILING DATE: 2004-11-03
;  PRIOR APPLICATION NUMBER: EP 04105507.0
;  PRIOR FILING DATE: 2004-11-03
;  PRIOR APPLICATION NUMBER: EP 04105485.9
;  PRIOR FILING DATE: 2004-11-03
;  PRIOR APPLICATION NUMBER: EP 04105484.2
;  PRIOR FILING DATE: 2004-11-03
;  PRIOR APPLICATION NUMBER: US 60/662,276
;  PRIOR FILING DATE: 2005-03-14
;  PRIOR APPLICATION NUMBER: US 60/700,293
;  PRIOR FILING DATE: 2005-07-18
;  NUMBER OF SEQ ID NOS: 483996
;  SOFTWARE: PatentIn version 3.3
; SEQ ID NO 227410
;  LENGTH: 1313
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;    TYPE: DNA
;    ORGANISM: Homo Sapiens
US-11-266-748A-227410

    Query Match            61.5%;  Score 1288;  DB 8;  Length 1313;
    Best Local Similarity  99.9%;  Pred. No. 2e-194;
    Matches 1299; Conservative    0; Mismatches    0; Indels    1; Gaps    1;

Qy          1 CTCCTCTGCTCCTCGAAGAAGGCCAGGGCGGGGCTGCCGCAAGTTTTGACATTTTCGCAG 60
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Db         15 CTCCTCTGCTCCTCGAAGAAGGCCAGGGCGGGGCTGCCGCAAGTTTTGACATTTTCGCAG 74

Qy         61 CGGAGACGCGCGCGGGCACTCTCGGGCCGACGGCTGCGGCGGCGGCCGACCCTCCAGAGC 120
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         75 CGGAGACGCGCGCGGGCACTCTCGGGCCGACGGCTGCGGCGGCGGCCGACCCTCCAGAGC 134

Qy        121 CCCTTAGTCGCGCCCCGGCCCTCCCGCTGCCCGGAGTCCGGCGGCCACGAGGCCCAGCCG 180
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy        181 CGTCCTCCCGCGCTTGCTCGCCCGGCGGCCGCAGCCATGTCCCGGGGGCCCGAGGAGGTG 240
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Db        195 CGTCCTCCCGCGCTTGCTCGCCCGGCGGCCGCAGCCATGTCCCGGGGGCCCGAGGAGGTG 254

Qy        241 AACCGGCTCACGGAGAGCACCTACCGGAATGTTATGGAACAGTTCAATCCTGGGCTGCGA 300
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Db        315 AATTTAATAAACCTGGGGAAAAATTATGAGAAAGCTGTAAACGCTATGATCCTGGCAGGA 374

Qy        361 AAAGCCTACTACGATGGAGTGGCCAAGATCGGTGAGATTGCCACTGGGTCCCCCGTGTCA 420
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Qy        421 ACTGAACTGGGACATGTCCTCATAGAGATTTCAAGTACCCACAAGAAACTCAACGAGAGT 480
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Db        435 ACTGAACTGGGACATGTCCTCATAGAGATTTCAAGTACCCACAAGAAACTCAACGAGAGT 494

Qy        481 CTTGATGAAAATTTTAAAAAATTCCACAAAGAGATTATCCATGAGCTGGAGAAGAAGATA 540
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Qy        601 AAAATTAGAGTCTTTGGAGAAATCCCAAGCTGAGTTGAAGAAGATCAGAAGGAAAAGCCAA 660
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Qy   1021 GTGGTTAGGAAAGATTACGACACCCTTTCTAAATGCTCACCAAAGATGCCCCCCGCTCCT 1080
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Qy   1081 TCAGGCAGAGCATATACCAGTCCCTTGATCGATATGTTTAATAACCCAGCCACGGCTGCC 1140
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Qy   1141 CCGAATTCACAAAGGGTAAATAATTCAACAGGTACTTCCGAAGATCCCAGTTTACAGCGA 1200
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Qy   1201 TCAGTTTCGGTTGCAACGGGACTGAACATGATGAAGAAGCAGAAAGTGAAGACCATCTTC 1260
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RESULT 2
US-11-266-748A-289813
; Sequence 289813, Application US/11266748A
; Publication No. US20060134663A1
; GENERAL INFORMATION:
;   APPLICANT: Harkin, Paul
;   APPLICANT:  Johnston, Patrick
;   APPLICANT:  Mulligan, Karl
;   TITLE OF INVENTION: Transcriptome Microarray Technology and
;   TITLE OF INVENTION:  Methods of Using the Same
;   FILE REFERENCE: 55815-0102 (319189)
;   CURRENT APPLICATION NUMBER: US/11/266,748A
;   CURRENT FILING DATE:  2005-11-03
;   PRIOR APPLICATION NUMBER: EP 04105479.2
;   PRIOR FILING DATE: 2004-11-03
;   PRIOR APPLICATION NUMBER: EP 04105482.6
;   PRIOR FILING DATE: 2004-11-03
;   PRIOR APPLICATION NUMBER: EP 04105483.4
;   PRIOR FILING DATE: 2004-11-03
;   PRIOR APPLICATION NUMBER: EP 04105507.0
;   PRIOR FILING DATE: 2004-11-03
;   PRIOR APPLICATION NUMBER: EP 04105485.9
;   PRIOR FILING DATE: 2004-11-03
;   PRIOR APPLICATION NUMBER: EP 04105484.2
;   PRIOR FILING DATE: 2004-11-03
;   PRIOR APPLICATION NUMBER: US 60/662,276
;   PRIOR FILING DATE: 2005-03-14
;   PRIOR APPLICATION NUMBER: US 60/700,293
;   PRIOR FILING DATE: 2005-07-18
;   NUMBER OF SEQ ID NOS: 483996
;   SOFTWARE: PatentIn version 3.3
; SEQ ID NO 289813
.;   LENGTH: 1000
;   TYPE: DNA
;   ORGANISM: Homo Sapiens
US-11-266-748A-289813

   Query Match         47.7%;  Score 1000;  DB 8;  Length 1000;
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   Matches 1000;  Conservative    0;  Mismatches   0;  Indels    0;  Gaps    0;

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RESULT 3
US-11-266-748A-341242/c
; Sequence 341242, Application US/11266748A
; Publication No. US20060134663A1
; GENERAL INFORMATION:
;  APPLICANT: Harkin, Paul
;  APPLICANT:  Johnston, Patrick
;  APPLICANT:  Mulligan, Karl
;  TITLE OF INVENTION: Transcriptome Microarray Technology and
;  TITLE OF INVENTION:  Methods of Using the Same
;  FILE REFERENCE: 55815-0102 (319189)
;  CURRENT APPLICATION NUMBER: US/11/266,748A
;  CURRENT FILING DATE:  2005-11-03
;  PRIOR APPLICATION NUMBER: EP 04105479.2
;  PRIOR FILING DATE: 2004-11-03
;  PRIOR APPLICATION NUMBER: EP 04105482.6
;  PRIOR FILING DATE: 2004-11-03
;  PRIOR APPLICATION NUMBER: EP 04105483.4
;  PRIOR FILING DATE: 2004-11-03
;  PRIOR APPLICATION NUMBER: EP 04105507.0
;  PRIOR FILING DATE: 2004-11-03
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